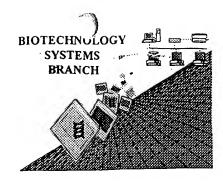
RAW SEQUENCE LISTING ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09/807,660
Source:	PCTO9
	5-1-01
Date Processed by STIC:	

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 c-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 c-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER</u> <u>VERSION 3.0 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 – 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address: http://www.uspto.gov/web/offices/pac/checker

PCT

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DATE: 05/01/2001
                     RAW SEQUENCE LISTING
                    PATENT APPLICATION: US/09/807,660
                                                           TIME: 11:37:50
                                                                               Does Not Comply
                     Input Set : A:\23,495 PCT.txt
                                                                           Corrected Diskette Needed
                    Output Set: N:\CRF3\05012001\I807660.raw
                                                                             see pp. 1,2
    3 <110> APPLICANT: Maliszewski, Charles R.
              Gayle III, Richard B.
              Marcus, Aaron J.
              Immunex Corporation
              Cornell Research Foundation, Inc.
      9 <120> TITLE OF INVENTION: Methods of Inhibiting Platelet Activation and
             Recruitment
     12 <130> FILE REFERENCE: 23,495 PCT
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/807,660
C--> 15 <141> CURRENT FILING DATE: 2001-04-16
     17 <150> PRIOR APPLICATION NUMBER: US 60/104,585
     18 <151> PRIOR FILING DATE: 1998-10-16
     20 <150> PRIOR APPLICATION NUMBER: US 60/107,466
     21 <151> PRIOR FILING DATE: 1998-11-06
     23 <150> PRIOR APPLICATION NUMBER: US 60/149,010
     24 <151> PRIOR FILING DATE: 1999-08-13
     26 <160> NUMBER OF SEQ ID NOS: 31
     28 <170> SOFTWARE: PatentIn Ver. 2.0
ERRORED SEQUENCES
     1925 <210> SEQ ID NO: 31
     1926 <211> LENGTH: 58
     1927 <212> TYPE: PRT
     1928 <213> ORGANISM: Homo sapiens
     1930 <400> SEQUENCE: 31
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                                              10
     1934 Val Cys Ser Ala Val Ser His Arg Asn Gln Gln Thr Trp Phe Glu Gly
                                          25
     1937 Ile Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Ser Thr Leu
                  35
     1940 Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr
     1941
E--> 1945 1
E--> 1948 1
              > Delete extroneous numerals at
                   the end of the file. They are causing an invalid amino acid count for sequence #31.
```

Seg #6 (Partially shown)
 Seg #6 (Partially shown)
 Assume the segmence of the sequence of the s

VERIFICATION SUMMARY DATE: 05/01/2001 PATENT APPLICATION: US/09/807,660 TIME: 11:37:52

Input Set : A:\23,495 PCT.txt

Output Set: N:\CRF3\05012001\I807660.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 L:620 M:258 W: Mandatory Feature missing, <220> FEATURE: L:620 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: L:850 M:258 W: Mandatory Feature missing, <220> FEATURE: L:850 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:

L:1426 M:258 W: Mandatory Feature missing, <220> FEATURE:

L:1426 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION: L:1945 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:31

M:332 Repeated in SeqNo=31